Docket No.: 1254-0258PUS1

**AMENDMENTS TO THE CLAIMS** 

This listing of claims will replace all prior versions, and listings, of claims in the present

application:

Listing of Claims:

1. (Currently Amended) [[A]] An isolated or purified gene coding for either one of the

following proteins (a) or (b):

(a) a protein consisting of the amino acid sequence shown in SEQ ID NO:2; or

(b) a protein consisting of an amino acid sequence shown in SEQ ID NO:2 wherein said

amino acid sequence is obtained by deletion, substitution or addition of one or more amino

[[acids,]] acids in SEQ. ID. NO.:2, wherein the amino acid at position 75 is methionine, and

wherein said protein has which exhibits scytalone dehydratase activity in the presence of a

scytalone dehydratase inhibitor.

2. (Currently Amended) [[A]] The gene according to claim 1, wherein the scytalone

dehydratase inhibitor inhibits the dehydration reaction from scytalone to 1,3,8-

trihydroxynaphtalene in a melanin biosynthesis pathway.

3. (Currently Amended) [[A]] The gene according to claim 1, wherein the scytalone

dehydratase inhibitor is carpropamid.

4. (Withdrawn) A scytalone dehydratase encoded by the gene of claim 1.

3

**GMM/ETP** 

Application No. 10/507,132 Art Unit 1652 Reply to Office Action of February 12, 2007

5. (Currently Amended) A recombinant vector comprising the gene of claim 1. a gene coding for either one of the following proteins (a) or (b):

(a) a protein consisting of the amino acid sequence shown in SEQ ID NO:2; or

(b) a protein consisting of an amino acid sequence wherein said amino acid sequence is obtained by deletion, substitution or addition of one or more amino acids in SEQ. ID. NO.:2, wherein the amino acid at position 75 is methionine, and wherein said protein has scytalone dehydratase activity in the presence of a scytalone dehydratase inhibitor.

6. (Currently Amended) A transformant obtained by transformation [[of]] with the recombinant vector of claim 5.

- 7. (Withdrawn) A method for assessing sensitivity of a rice blast fungus to a scytalone dehydratase inhibitor, comprising the steps of:
- (a) identifying an amino acid in an amino acid sequence of scytalone dehydratase in a subject rice blast fungus, which corresponds to valine at position 75 in the amino acid sequence shown in SEQ ID NO: 4; and
- (b) assessing sensitivity of the subject rice blast fungus to the scytalone dehydratase inhibitor based on the results of step (a).
- 8. (Withdrawn) A method for assessing sensitivity according to claim 7, wherein when the amino acid identified in step (a) is methionine, the sensitivity of the subject rice blast fungus

Docket No.: 1254-0258PUS1

Application No. 10/507,132

Art Unit 1652

Reply to Office Action of February 12, 2007

to the scytalone dehydratase inhibitor is assessed to be lower than that of a wild-type rice blast fungus in step (b).

9. (Withdrawn) A kit for screening an inhibitor, comprising the scytalone dehydratase of claim 4.

10. (Currently Amended) A kit for assessing a rice blast fungus resistant to a scytalone dehydratase inhibitor, said kit comprising a pair of primers designed to flank a nucleotide sequence coding for an amino acid corresponding to valine at position 75 in the amino acid sequence shown in SEQ ID NO: 4.

- 11. (Currently Amended) A kit for assessing a rice blast fungus resistant to a scytalone dehydratase inhibitor, said kit comprising an oligonucleotide including a nucleotide sequence coding for an amino acid corresponding to valine at position 75 in the amino acid sequence shown in SEQ ID NO: 4.
- 12. (New) An isolated or purified gene coding for a protein comprising of the amino acid sequence shown in SEQ ID NO:2.
- 13. (New) The isolated or purified gene of claim 12, wherein the protein consists of SEQ ID NO:2.

5 GMM/ETP

14. (New) The gene according to claim 1, wherein the gene codes for protein (b) and

Docket No.: 1254-0258PUS1

hybridizes to a nucleotide sequence complementary to the nucleotide sequence of SEQ. ID.

NO.:1 under stringent conditions.

15. (New) The gene according to claim 14, wherein the stringent conditions include a

sodium concentration of 10-300 mM and a temperature of 25-70°C.

16. (New) The gene according to claim 14, wherein the stringent conditions include a

sodium concentration of 20-100 mM and a temperature of 42-55°C.

17. (New) A recombinant vector comprising a gene coding for a protein comprising of

the amino acid sequence shown in SEQ ID NO:2.

18. (New) A transformant obtained by transformation with the recombinant vector of

claim 17.

19. (New) An isolated or purified gene coding for a protein consisting of an amino acid

sequence wherein said amino acid sequence is obtained by deletion, substitution or addition of 1-

30 amino acids in SEQ. ID. NO.:2, wherein the amino acid at position 75 is methionine, and

wherein said protein has scytalone dehydratase activity in the presence of a scytalone

dehydratase inhibitor.

6 GMM/ETP

Application No. 10/507,132 Art Unit 1652 Reply to Office Action of February 12, 2007 Docket No.: 1254-0258PUS1

- 20. (New) The gene of claim 19, which contains 1-20 of said deletions, substitutions or additions.
- 21. (New) The gene of claim 19, which contains 1-10 of said deletions, substitutions or additions.